

SEQUENCE LISTING

<110> Bayer Aktiengesellschaft

<120> Ultraspiracle (USP) protein from *Heliothis virescens*

<130> Le A 34 771

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<150> DE 100 36 469.1

<151> 2000-07-25

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<170> PatentIn Ver. 2.1

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<212> DNA

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acg cct acg tcg ccc gga aac atg ctt caa ccg atg gct acg ccg tct     144
Thr Pro Thr Ser Pro Gly Asn Met Leu Gln Pro Met Ala Thr Pro Ser
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aac tta ccg act gtc gac tgc tca ctc gat att caa tgg cta aac ttg     192
Asn Leu Pro Thr Val Asp Cys Ser Leu Asp Ile Gln Trp Leu Asn Leu
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gag gga ggt ttt atg tcg ccg atg tca ccg ccg gag atg aag cca gac     240
Glu Gly Gly Phe Met Ser Pro Met Ser Pro Pro Glu Met Lys Pro Asp
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ttc aag aac tac ccc ccg aac cat ccc cta agt ggt tct aag cac ctc 336
 Phe Lys Asn Tyr Pro Pro Asn His Pro Leu Ser Gly Ser Lys His Leu
 100 105 110

tgt tct ata tgt gga gat aga gcg tcg ggg aaa cat tat gga gta tac 384
 Cys Ser Ile Cys Gly Asp Arg Ala Ser Gly Lys His Tyr Gly Val Tyr
 115 120 125

agt tgt gaa ggt tgc aaa ggt ttc ttc aaa agg acg gta aga aaa gac 432
 Ser Cys Glu Gly Cys Lys Gly Phe Phe Lys Arg Thr Val Arg Lys Asp
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tca atc gag cgg ttg ctg gag atg gag tca ctg gta gct gac ccc agc 672
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Asn Leu Pro Thr Val Asp Cys Ser Leu Asp Ile Gln Trp Leu Asn Leu
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Glu Gly Gly Phe Met Ser Pro Met Ser Pro Pro Glu Met Lys Pro Asp
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Thr Ala Met Leu Asp Gly Leu Arg Asp Asp Ser Thr Pro Pro Pro Ala
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Phe Lys Asn Tyr Pro Pro Asn His Pro Leu Ser Gly Ser Lys His Leu
100 105 110

Cys Ser Ile Cys Gly Asp Arg Ala Ser Gly Lys His Tyr Gly Val Tyr
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Ser Cys Glu Gly Cys Lys Gly Phe Phe Lys Arg Thr Val Arg Lys Asp
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Leu Thr Tyr Ala Cys Arg Glu Glu Arg Asn Cys Ile Ile Asp Lys Arg
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Gln Arg Asn Arg Cys Gln Tyr Cys Arg Tyr Gln Lys Cys Leu Ala Cys
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Gly Met Lys Arg Glu Ala Val Gln Glu Glu Arg Gln Arg Ala Ala Arg
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Gly Thr Glu Asp Ala His Pro Ser Ser Ser Val Gln Val Gln Glu Leu

195

200

205

Ser Ile Glu Arg Leu Leu Glu Met Glu Ser Leu Val Ala Asp Pro Ser
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Lys Phe Arg Ala Pro Val Ser Ser Leu Cys Gln Ile Gly Asn Lys Gln
245 250 255

Ile Ala Ala Leu Val Val Trp Ala Arg Asp Ile Pro His Phe Ser Gln
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Leu Glu Met Glu Asp Gln Ile Leu Leu Ile Lys Gly Ser Trp Asn Glu
275 280 285

Leu Leu Leu Phe Ala Ile Ala Trp Arg Ser Met Glu Phe Leu Thr Glu
290 295 300

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305 310 315 320

Gln Leu Met Cys Leu Met Pro Gly Met Thr Leu His Arg Asn Ser Ala
325 330 335

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Ser Leu Lys Met Arg Thr Leu Arg Val Asp Gln Ala Glu Tyr Val Ala
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Arg Gln Glu Val Glu Val Leu Arg Glu Lys Met Phe Leu Cys Leu Asp
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Leu Leu Leu Arg Leu Pro Ala Leu Arg Ser Ile Ser Leu Lys Ser Phe
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Tyr Ile Arg Asp Ala Leu Arg Asn His Ala Pro Pro Ile Asp Thr Asn

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